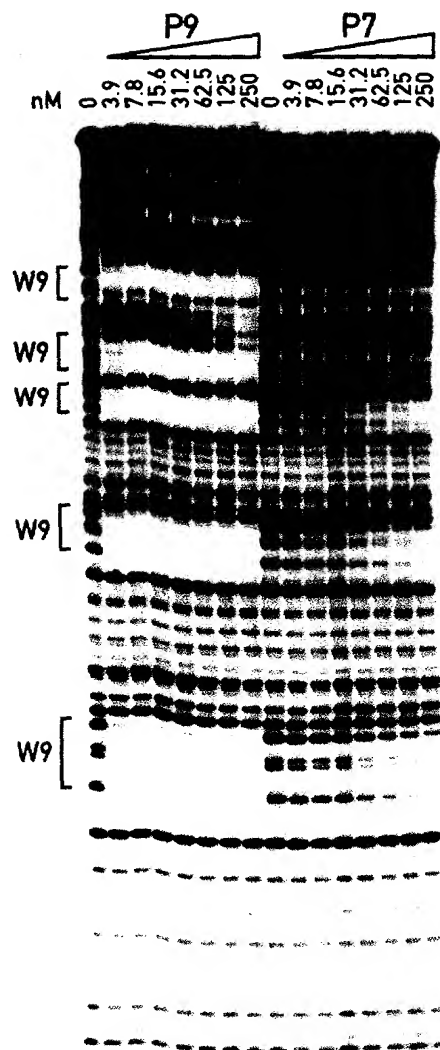


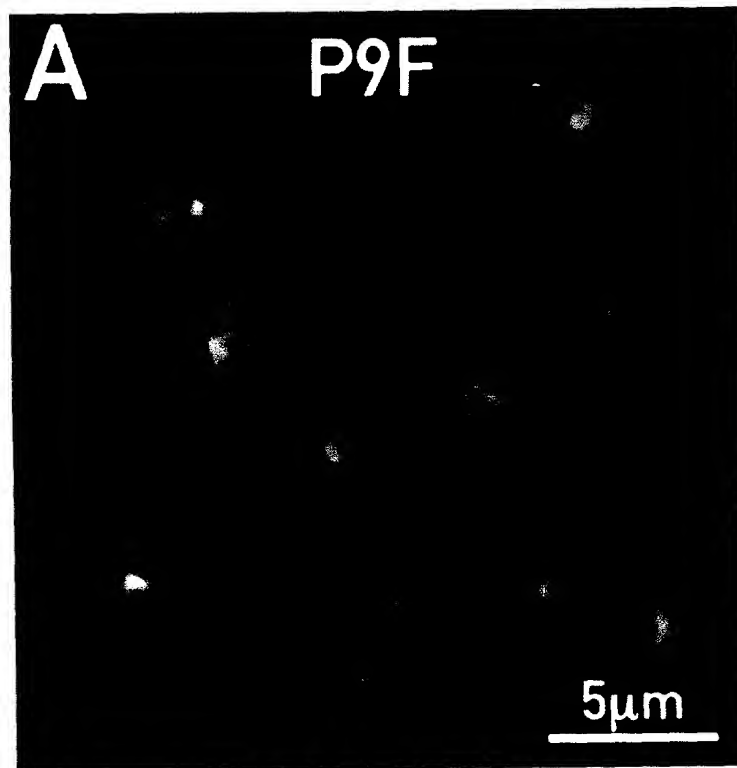
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FIGURE 1



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FIGURE 2



A

nM P31
0 0.1 0.25 1 5 25
P31E

G 0 0.25 1 5 0 P31E

5'-CGTAATCATGGTTCATAGC-3'

5'-AACATAAAAGTGTAAGCG-3'

Ubx

(AAGAG)₂

(AAGAG)₅

(AACAC)₂

(AAGAG)₅

(AACAC)₂

(AAGAG)₅

(AACAC)₂

1 2 3 4 5 6 7 8 9 10 11 12 13 14 15 16

GAF31 Brown I

B

GAGA factor (GAF)

0 1 2 3 4 5 6 7 8 9 10 11 12 13

Ubx

(AAGAG)₂

(AAGAG)₅

(AACAC)₂

(AAGAG)₅

(AACAC)₂

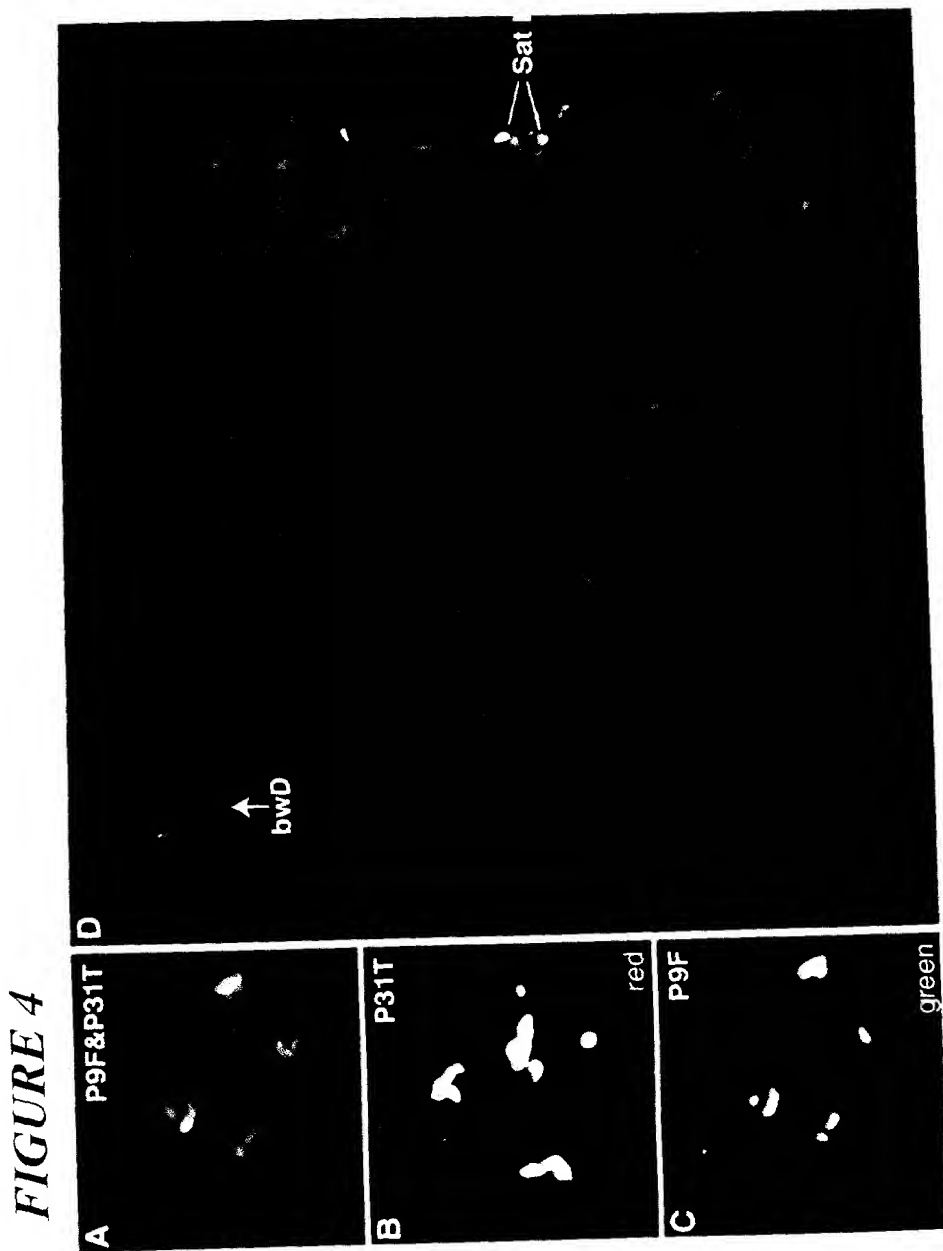
(AAGAG)₅

(AACAC)₂

1 2 3 4 5 6 7 8 9 10 11 12 13

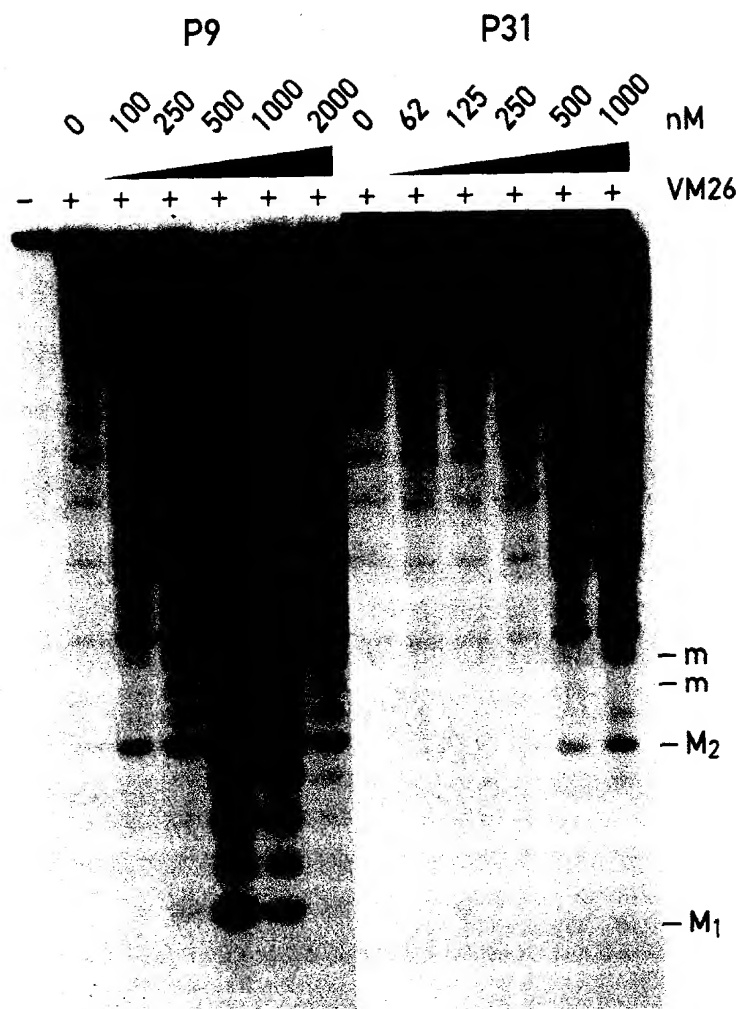
GAF31

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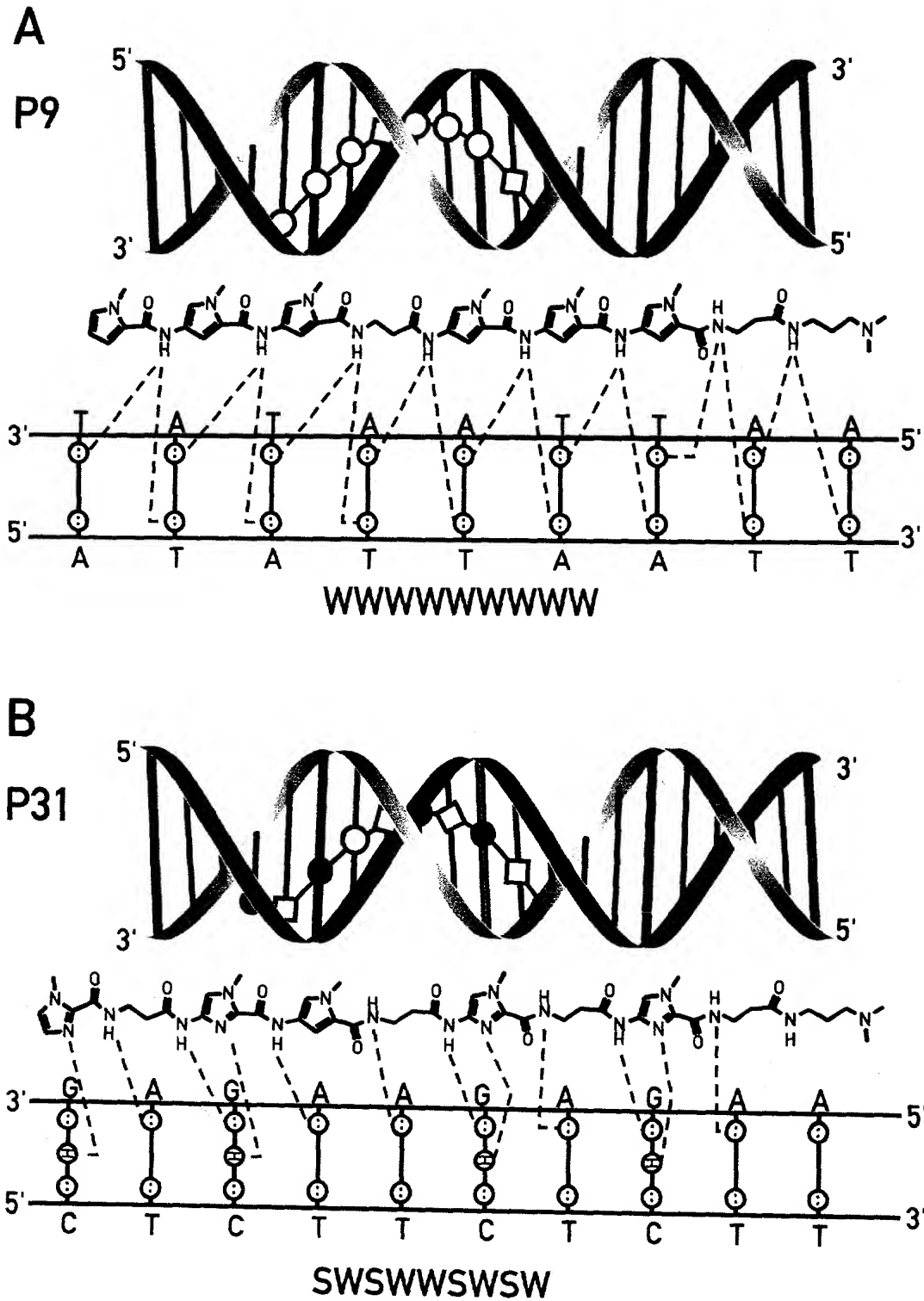
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FIGURE 5



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FIGURE 6

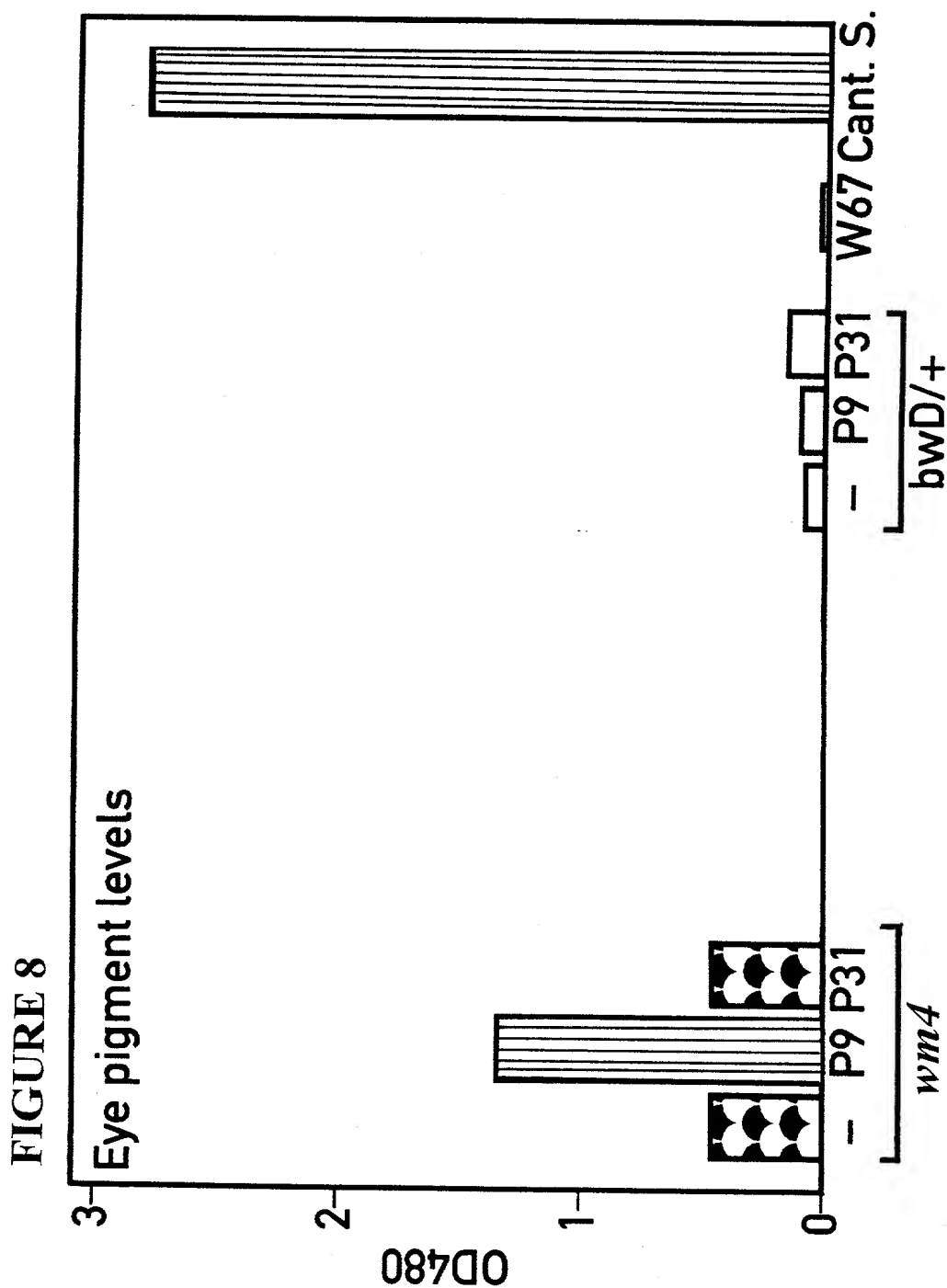


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FIGURE 7

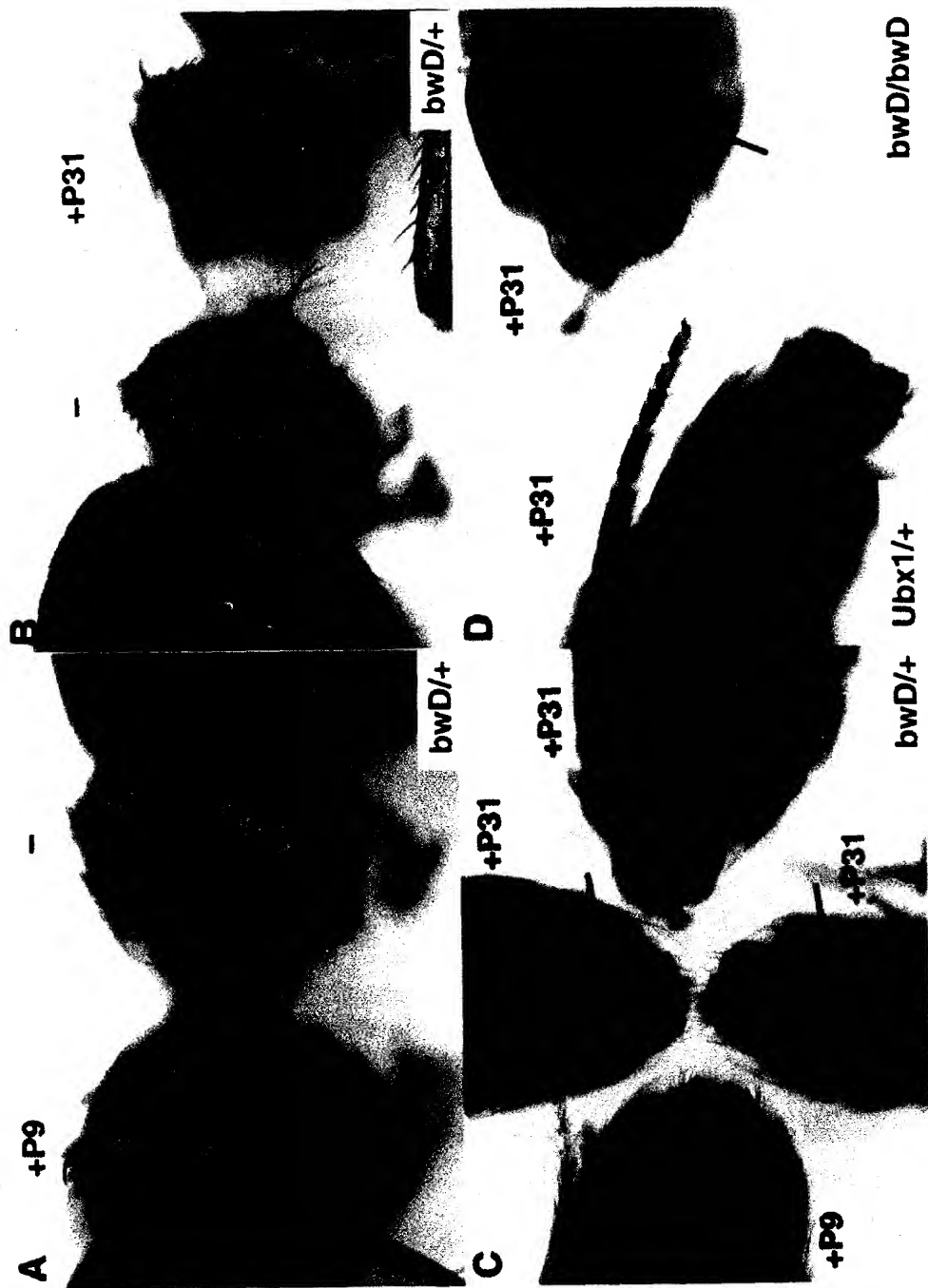


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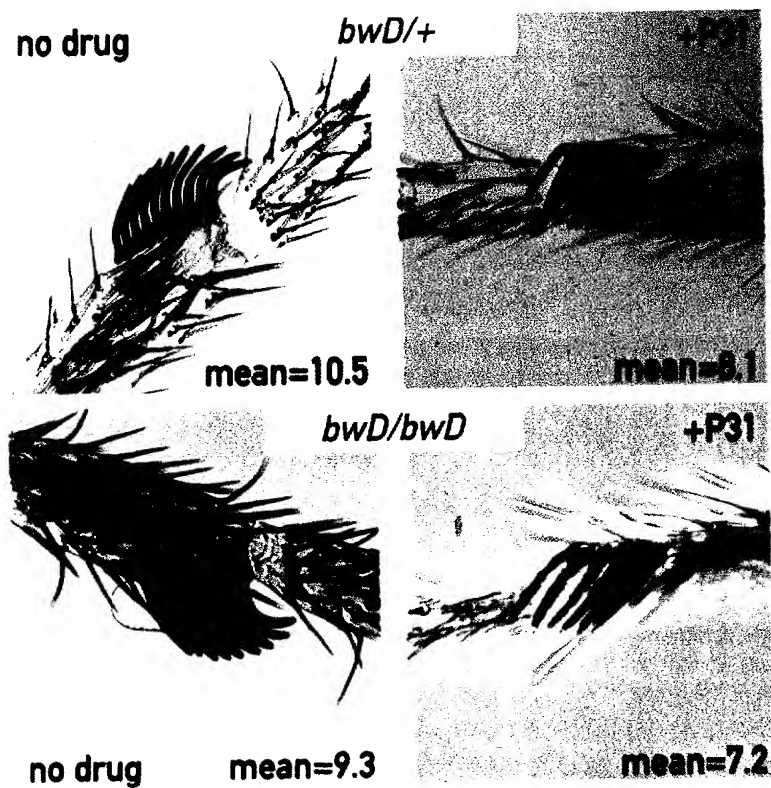
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FIGURE 9



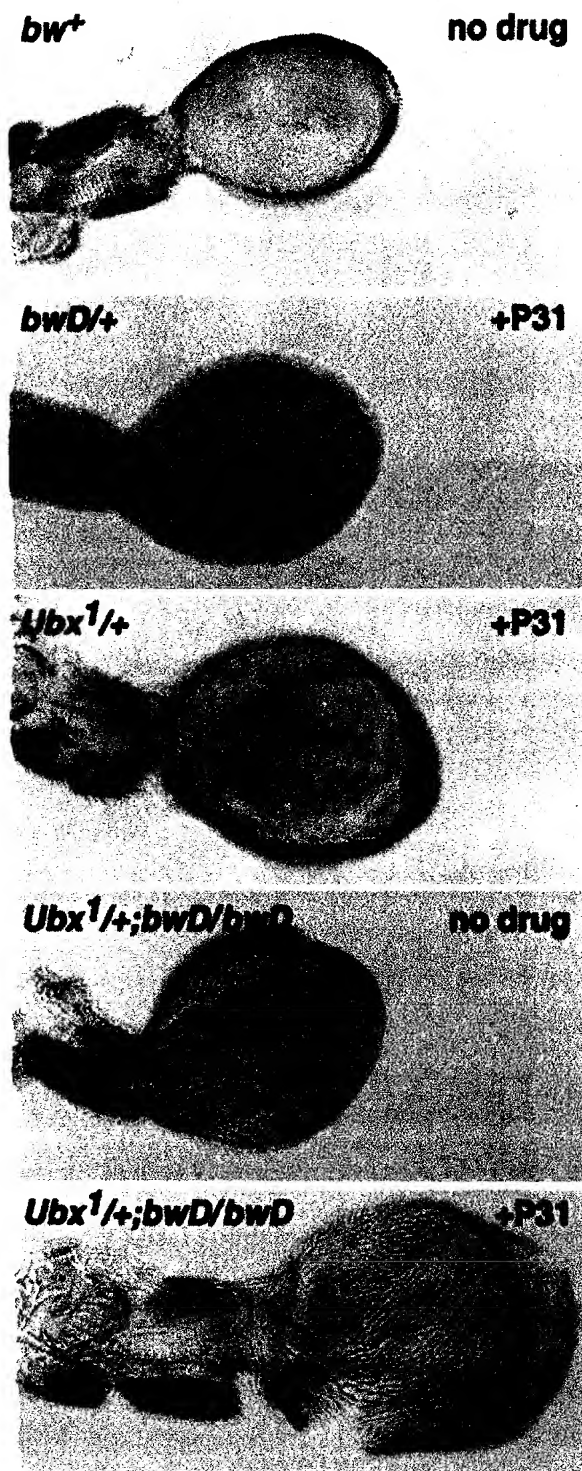
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FIGURE 10



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FIGURE 11



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FIGURE 12

